## SEQUENCE LISTING

<110>	Holloway, James L.
<120>	Human Serine Protease
<130>	99-88C1
	60/167,038 1999-11-23
	09/715,994 2000-11-17
<160>	4
<170>	FastSEQ for Windows Version 3.0
<210> <211> <212> <213>	807
<220> <221> <222>	
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	gtg gtg gag gac act tcg aga gtg cca aat gac aag tgg 96 Val Val Glu Asp Thr Ser Arg Val Pro Asn Asp Lys Trp 20 25 30
	ggg gcc ctg ctc tct gcg tcc tgg atc ctc aca gca gct Gly Ala Leu Leu Ser Ala Ser Trp Ile Leu Thr Ala Ala 40 45
	cgc tcc cag cgt aga gac acc acg gtg ata cca gtc tcc Arg Ser Gln Arg Arg Asp Thr Thr Val Ile Pro Val Ser 55 60

_		cat His	_		~		_		_		-		-	-		240
_		gca Ala	-													288
		atc Ile						_		-	-		_	_	_	336
		gtg Val 115		_				_								384
		cct Pro	-		_	_			-	-						432
		atc Ile								_						480
		cgg Arg		_		-	-	-	-							528
		cac His	-													576
	_	gtc Val 195	_			_		_	_							624
		acg Thr														672
gac	ttg	agc	cag	cgc	tgg	gtg	gtg	caa	ggc	ctg	gtg	tcc	tgg	<b>ggg</b>	gga	720

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Asp Leu Ser Gln Arg Trp Val Val Gln Gly Leu Val Ser Trp Gly Gly
225
                                                                      768
cct gaa gaa tgc ggc agc aag cag gtc tat gga gtc tac aca aag gtc
Pro Glu Glu Cys Gly Ser Lys Gln Val Tyr Gly Val Tyr Thr Lys Val
                                    250
                                                         255
                245
                                                                      807
tcc aat tac gtg gac tgg gtg tgg gag cag atg ggc tta
Ser Asn Tyr Val Asp Trp Val Trp Glu Gln Met Gly Leu
            260
      <210> 2
      <211> 269
      <212> PRT
      <213> Homo sapiens
      <400> 2
Arg Ile Ile Gly Gly Arg Asn Ala Glu Pro Gly Leu Phe Pro Trp Gln
                 5
                                    10
Ala Leu Ile Val Val Glu Asp Thr Ser Arg Val Pro Asn Asp Lys Trp
Phe Gly Ser Gly Ala Leu Leu Ser Ala Ser Trp Ile Leu Thr Ala Ala
His Val Leu Arg Ser Gln Arg Arg Asp Thr Thr Val Ile Pro Val Ser
Lys Glu His Val Thr Val Tyr Leu Gly Leu His Asp Val Arg Asp Lys
Ser Gly Ala Val Asn Ser Ser Ala Ala Arg Val Val Leu His Pro Asp
                                    90
                85
Phe Asn Ile Gln Asn Tyr Asn His Asp Ile Ala Leu Val Gln Leu Gln
            1.00
                                105
                                                     110
Glu Pro Val Pro Leu Gly Pro His Val Met Pro Val Cys Leu Pro Arg
                            120
Leu Glu Pro Glu Gly Pro Ala Pro His Met Leu Gly Leu Val Ala Gly
                        135
                                             140
Trp Gly Ile Ser Asn Pro Asn Val Thr Val Asp Glu Ile Ile Ser Ser
145
                    150
                                         155
Gly Thr Arg Thr Leu Ser Asp Val Leu Gln Tyr Val Lys Leu Pro Val
                165
                                    170
Val Pro His Ala Glu Cys Lys Thr Ser Tyr Glu Ser Arg Ser Gly Asn
            180
                                185
Tyr Ser Val Thr Glu Asn Met Phe Cys Ala Gly Tyr Tyr Glu Gly Gly
        195
                            200
                                                 205
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Lys Asp Thr Cys Leu Gly Asp Ser Gly Gly Ala Phe Val Ile Phe Asp
    210
                        215
                                            220
Asp Leu Ser Gln Arg Trp Val Val Gln Gly Leu Val Ser Trp Gly Gly
225
                    230
                                        235
Pro Glu Glu Cys Gly Ser Lys Gln Val Tyr Gly Val Tyr Thr Lys Val
                                    250
                                                         255
                245
Ser Asn Tyr Val Asp Trp Val Trp Glu Gln Met Gly Leu
                                265
            260
      <210> 3
      <211> 807
      <212> DNA
      <213> Artificial Seguence
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      <223> This degenerate sequence encodes the amino acid
            sequence of SEQ ID NO:2.
      <221> variation
      <222> (1)...(807)
      <223> N is any nucleotide.
      <400> 3
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gtngargaya cnwsnmgngt nccnaaygay aartggttyg gnwsnggngc nytnytnwsn
gcnwsntgga thytnacngc ngcncaygtn ytnmgnwsnc armgnmgnga yacnacngtn
athcongtnw snaargarca ygtnacngtn tayytnggny tncaygaygt nmgngayaar
wsnggngcng tnaaywsnws ngengenmgn gtngtnytne ayeengaytt yaayathear
aaytayaayc aygayathgc nytngtncar ytncargarc cngtnccnyt nggnccncay
qtnatgccnq tntqyytncc nmgnytngar ccngarggnc cngcnccnca yatgytnggn
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ytngtngcng gntggggnat hwsnaayccn aaygtnacng tngaygarat hathwsnwsn

ggnacnmgna cnytnwsnga ygtnytncar taygtnaary tnccngtngt nccncaygcn gartgyaara cnwsntayga rwsnmgnwsn ggnaaytayw sngtnacnga raayatgtty

tgygcnggnt aytayqarqg nggnaargay acntgyytng gngaywsngg nggngcntty

qtnathttyq aygayytnws ncarmqntqq qtnqtncarq qnytnqtnws ntqqqqnqqn

ccngargart gyggnwsnaa rcargtntay ggngtntaya cnaargtnws naaytaygtn

60

120

180

240

300

360 420

480 540

600

660

720

780

807

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<212> PRT

gaytgggtnt gggarcarat gggnytn

<213> Artificial Sequence

<220> <223> Peptide linker.